

## Appendix A

>Top sequence C5aR human 350 aa vs.  
>Bottom sequence C3aR mouse 347 aa

65.6% identity overall  
N-terminal and 2<sup>nd</sup> extracellular loop are underlined, 29.0% identity

	10	20	30	40	50
C5aR human	<u>MNSFN</u> <u>YTPDYGHYDDK</u> <u>DTLDL</u> <u>NTFPV</u> <u>DKTSNTLR</u> <u>VP</u> -DILALVIFAVVFLVGLNALVV				
	:::..	:::..	:::..	:::..	:::..
C5aR mouse	MNS-GFEI-NYDHYG---TMDPNIPADGIHLPKRQPGDVAALIITYSVVFLVGVPGNALVV				
	10	20	30	40	50
	60	70	80	90	100
C5aR human	WVTAFEAKRTINAIWFLNLAVADFLSCLALPILFTSIVQHHHWPFGGAACSIPLSLILLN				
	:::..	:::..	:::..	:::..	:::..
C5aR mouse	WVTAFEPDGPSNAIWFLNLAVADLLSCLAMPVLFITVLNHNHYWYFDATAIVLPSLILLN				
	60	70	80	90	100
	120	130	140	150	160
C5aR human	MYASILLLATISADRFLLVFKPIWCQNFAGLAWIACAVAWGLALLLTIPSFY <u>RVVRE</u>				
	:::..	:::..	:::..	:::..	:::..
C5aR mouse	MYASILLLATISADRFLLVFKPIWCQKVRGTGLANMACGVAVWGLALLLTIPSFVYREAYK				
	120	130	140	150	160
	180	190	200	210	220
C5aR human	<u>EYFP</u> <u>PKVLCGV</u> <u>DYSHDKR</u> -RERAVAIVRLVLGFLWPLLTITICYTIFILLRTWSRRATRST				
	...	...	...	...	...
C5aR mouse	DFYSEHTVCGINYGGSFPKEKAVAILRLMWGFLVPLLTINICVTFLLLRTWSRKATRST				
	180	190	200	210	220
	240	250	260	270	280
C5aR human	KTLKVVAVVAVSFFIFWLPYQVTGIMMSFLEPSSPTFLLNKLDLCLVFSFAYINCCINPI				
	:::..	:::..	:::..	:::..	:::..
C5aR mouse	KTLKVVMAVVICFFIFWLPYQVTGVMIAWLPPSSPTLKRVEKLNSLCVSLAYINCCVNIPI				
	240	250	260	270	280
	300	310	320	330	340
C5aR human	IYVVGQGFQGLRKSLPSLLRNVLTEESVVRSEKSFTRSTVDTMAQKTQAV				
	:::..	:::..	:::..	:::..	:::..
C5aR mouse	IYVMAGQGFHGLRLSLPSIIRNALSEDSVGRDSKTFITPSTDDTSRKRSQAV				
	300	310	320	330	340